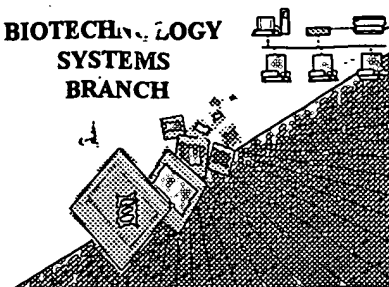


RAW SEQUENCE LISTING
ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



US 70
0905

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/938,275

Source: OIP

Date Processed by STIC: 9/14/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

OIPE

RAW SEQUENCE LISTING

DATE: 09/14/2001

PATENT APPLICATION: US/09/938,275

TIME: 11:04:48

Input Set : A:\P03seqid.txt

Output Set: N:\CRF3\09142001\I938275.raw

SEQUENCE LISTING

Does Not Comply
Corrected Diskette Needed

Please insure this mandatory heading is shown

3 (1) GENERAL INFORMATION:

C--> 4 (i) APPLICANT: Gerardo Castillo and Alan Snow

5 (ii) TITLE OF INVENTION: Therapeutic and Diagnostic Applications

6 of Laminin and Laminin-Derived Protein Fragments

7 (iii) NUMBER OF SEQUENCES: 11

8 (iv) CORRESPONDENCE ADDRESS:

9 (A) ADDRESSEE: Patrick M. Dwyer

10 (B) STREET: 1919 One Union Square, 600 University Street

11 (C) CITY: Seattle

12 (D) STATE: WA (Washington)

13 (E) COUNTRY: United States of America

14 (F) ZIP: 98101

15 (v) COMPUTER READABLE FORM:

16 (A) MEDIUM TYPE: Diskette - 3.50 inch, 1.44 Mb storage

17 (B) COMPUTER: IBM PC

18 (C) OPERATING SYSTEM: PC-DOS (Windows NT Version 4.0, '95)

19 (D) SOFTWARE: WordPerfect 5.2

20 (vi) CURRENT APPLICATION DATA:

C--> 21 (A) APPLICATION NUMBER: US/09/938,275

C--> 22 (B) FILING DATE: 22-Aug-2001

23 (C) CLASSIFICATION: U.S. Provisional Appl.

24 (vii) PRIOR APPLICATION DATA:

25 (A) APPLICATION NUMBER: 60/027,981

26 (B) FILING DATE: 08-October-1996

28 (viii) ATTORNEY/AGENT INFORMATION:

29 (A) NAME: Dwyer, Patrick M.

30 (B) REGISTRATION NUMBER: 32,411

31 (C) REFERENCE/DOCKET NUMBER: PROTEO.P03

32 (ix) TELECOMMUNICATION INFORMATION:

33 (A) TELEPHONE: (206) 343-7074

34 (B) TELEFAX: (206) 343-7085

ERRORED SEQUENCES

See next page

Per 1.822 of Sequence Rules
 a MAXIMUM of 16 amino acids
 = per line

09/938,275 2

insert hard
 return

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

| | | | | | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Leu | Gln | Val | Gln | Leu | Ser | Ile | Arg | Thr | Phe | Ala | Ser | Ser | Gly | Leu | Ile | Tyr | Tyr | Val | Ala |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | | | | | 20 |
| His | Gln | Asn | Gln | Met | Asp | Tyr | Ala | Thr | Leu | Gln | Leu | Gln | Glu | Gly | Arg | Leu | His | Phe | Met |
| | | | | 25 | | | | | 30 | | | | | 35 | | | | | 40 |
| Phe | Asp | Leu | Gly | Lys | Gly | Arg | Thr | Lys | Val | Ser | His | Pro | Ala | Leu | Leu | Ser | Asp | Gly | Lys |
| | | | | 45 | | | | | 50 | | | | | 55 | | | | | 60 |
| Trp | His | Thr | Val | Lys | Thr | Glu | Tyr | Ile | Lys | Arg | Lys | Ala | Phe | Met | Thr | Val | Asp | Gly | Gln |
| | | | | 65 | | | | | 70 | | | | | 75 | | | | | 80 |
| Glu | Ser | Pro | Ser | Val | Thr | Val | Val | Gly | Asn | Ala | Thr | Thr | Leu | Asp | Val | Glu | Arg | Lys | Leu |
| | | | | 85 | | | | | 90 | | | | | 95 | | | | | 100 |
| Tyr | Leu | Gly | Gly | Leu | Pro | Ser | His | Tyr | Arg | Ala | Arg | Asn | Ile | Gly | Thr | Ile | Thr | His | Ser |
| | | | | 105 | | | | | 110 | | | | | 115 | | | | | 120 |
| Ile | Pro | Ala | Cys | Ile | Gly | Glu | Ile | Met | Val | Asn | Gly | Gln | Gln | Leu | Asp | Lys | Asp | Arg | Pro |
| | | | | 125 | | | | | 130 | | | | | 135 | | | | | 140 |
| Leu | Ser | Ala | Ser | Ala | Val | Asp | Arg | Cys | Tyr | Val | Val | Ala | Gln | Glu | Gly | Thr | Phe | Phe | Glu |
| | | | | 145 | | | | | 150 | | | | | 155 | | | | | 160 |
| Gly | Ser | Gly | Tyr | Ala | Ala | Leu | Val | Lys | Glu | Gly | Tyr | Lys | Val | Arg | Leu | Asp | | | |
| | | | | 165 | | | | | 170 | | | | | 175 | | | | | |

Please
 edit

↓

The types of errors shown exist throughout
 the Sequence Listing. Please check subsequent
 sequences for similar errors.